CRF Errors Corrected by the STIC Systems Branch

Serial Number: __09/965,594__ ENTERED

CRF Processing Date: __10/25/2001__
Edited by: _____
Verified by: ___AV___ (STIC sta)

☐ Changed a file from non-ASCII to ASCII.

☐ Changed the margins in cases where the sequence text was 'wrapped' down to the next line. #9788

☐ Edited a format error in the Current Application Data section, specifically: _____

☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____

☐ Added the mandatory heading and subheadings for 'Current Application Data'.

☐ Edited the 'Number of Sequences' field. The applicant spelled out a number instead of using an integer.

☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____

☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____

☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____

☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.

☐ Inserted colons after headings/subheadings. Headings edited included: _____

☐ Deleted extra, invalid, headings used by an applicant, specifically: _____

☑ Deleted: ☑ non-ASCII 'garbage' at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____

☐ Inserted mandatory headings, specifically: _____

☐ Corrected an obvious error in the response, specifically: _____

☐ Edited identifiers where upper case is used but lower case is required, or vice versa.

☐ Corrected an error in the Number of Sequences field, specifically: _____

☐ A 'Hard Page Break' code was inserted by the applicant. All occurrences had to be deleted.

☐ Deleted ending stop codon in amino acid sequences and adjusted the '(A)Length:' field accordingly (error due to a PatentIn bug). Sequences corrected: _____

☐ Other: _____
_____
_____
_____

Examiner: The above corrections must be communicated to the applicant in the first Office Action! DO NOT send a copy of this form.

3/1/95

#2

RAW SEQUENCE LISTING   DATE: 10/25/2001
PATENT APPLICATION: US/09/965,594   TIME: 08:46:14

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\10252001\I965594.raw

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  5 <110> APPLICANT: Wittekind, Michael
  7          Weinheimer, Steven
  9          Zhang, Yaqun
 11          Goldfarb, Valentina
 15 <120> TITLE OF INVENTION: Modified Forms of Hepatitis C NS3 Protease for
 17          Facilitating Inhibitor Screening and Structural Studies
 19          of Protease:Inhibitor Complexes
 23 <130> FILE REFERENCE: DB17Sequences
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C--> 27 <140> CURRENT APPLICATION NUMBER: US/09/965,594
C--> 29 <141> CURRENT FILING DATE: 2001-09-27
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 33 <150> PRIOR APPLICATION NUMBER: 60/115,271
 35 <151> PRIOR FILING DATE: 1999-01-08
 39 <160> NUMBER OF SEQ ID NOS: 26
 43 <170> SOFTWARE: PatentIn Ver. 2.0
 47 <210> SEQ ID NO: 1
 49 <211> LENGTH: 182
 51 <212> TYPE: PRT
 53 <213> ORGANISM: Hepatitis C virus
 57 <400> SEQUENCE: 1
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 65 Cys Ile Ile Thr Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly
 67             20                  25                  30
 71 Glu Val Gln Ile Val Ser Thr Ala Ala Gln Thr Phe Leu Ala Thr Cys
 73         35                  40                  45
 77 Ile Asn Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr
 79     50                  55                  60
 83 Ile Ala Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp
 85 65                  70                  75                  80
 89 Lys Asp Leu Val Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu Thr
 91             85                  90                  95
 95 Pro Cys Thr Cys Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala
 97             100                 105                 110
101 Asp Val Ile Pro Val Arg Arg Arg Gly Asp Ser Arg Gly Ser Leu Leu
103         115                 120                 125
107 Ser Pro Arg Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu
109         130                 135                 140
113 Leu Cys Pro Ala Gly His Ala Val Gly Ile Phe Arg Ala Ala Val Cys
115 145                 150                 155                 160
119 Thr Arg Gly Val Ala Lys Ala Val Asp Phe Ile Pro Val Glu Ser Leu
121                 165                 170                 175
125 Glu Thr Thr Met Arg Ser
127                 180
133 <210> SEQ ID NO: 2
135 <211> LENGTH: 549
137 <212> TYPE: DNA
139 <213> ORGANISM: Hepatitis C virus
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RAW SEQUENCE LISTING   DATE: 10/25/2001
PATENT APPLICATION: US/09/965,594   TIME: 08:46:14

Input Set : A:\PTO.AMC.txt
Output Set: N:\CRF3\10252001\I965594.raw

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149 gctcagacct tcctggctac ctgcatcaac ggtgtttgct ggaccgttta ccacggtgct 180
151 ggtacccgta ccatcgcttc cccgaaaggt ccggttatcc agatgtacac caacgttgac 240
153 aaagacctgg ttggttggcc ggctccgcag ggttcccgtt ccctgacccc gtgcacctgc 300
155 ggttcctccg acctgtacct ggttacccgt cacgctgacg ttatcccggt cgtcgtcgt 360
157 ggtgactccc gtggttccct gctgtccccg cgtccgatct cctacctgaa aggttcctcc 420
159 ggtggtccgc tgctgtgccc ggctggtcac gctgttggta tcttccgtgc tgctgtttgc 480
161 acccgtggtg ttgctaaagc tgttgacttc atcccggttg aatccctgga aaccaccatg 540
163 cgttcctga                                                        549
167 <210> SEQ ID NO: 3
169 <211> LENGTH: 195
171 <212> TYPE: PRT
173 <213> ORGANISM: Hepatitis C virus
177 <400> SEQUENCE: 3
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181  1               5                   10                  15
185 Gly Ala Tyr Ala Gln Gln Thr Arg Gly Leu Leu Gly Cys Ile Ile Thr
187             20                  25                  30
191 Ser Leu Thr Gly Arg Asp Lys Asn Gln Val Glu Gly Glu Val Gln Ile
193         35                  40                  45
197 Val Ser Thr Ala Ala Gln Thr Phe Leu Ala Thr Cys Ile Asn Gly Val
199     50                  55                  60
203 Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr Ile Ala Ser Pro
205 65                  70                  75                  80
209 Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp Lys Asp Leu Val
211             85                  90                  95
215 Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu Thr Pro Cys Thr Cys
217             100                 105                 110
221 Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala Asp Val Ile Pro
223         115                 120                 125
227 Val Arg Arg Arg Gly Asp Ser Arg Gly Ser Leu Leu Ser Pro Arg Pro
229     130                 135                 140
233 Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu Leu Cys Pro Ala
235 145                 150                 155                 160
239 Gly His Ala Val Gly Ile Phe Arg Ala Ala Val Cys Thr Arg Gly Val
241             165                 170                 175
245 Ala Lys Ala Val Asp Phe Ile Pro Val Glu Ser Leu Glu Thr Thr Met
247             180                 185                 190
251 Arg Ser Pro
253         195
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261 <211> LENGTH: 588
263 <212> TYPE: DNA
265 <213> ORGANISM: Hepatitis C virus
269 <400> SEQUENCE: 4
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273 cagcagactc gaggtctgct gggttgcatc atcacctccc tgaccggtcg tgacaaaaac 120
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275 caggttgaag gtgaagttca gatcgtttcc accgctgctc agaccttcct ggctacctgc 180
277 atcaacggtg tttgctggac cgtttaccac ggtgctggta cccgtaccat cgcttccccg 240
279 aaaggtccgg ttatccagat gtacaccaac gttgacaaag acctggttgg ttggccggct 300
281 ccgcagggtt cccgttccct gaccccgtgc acctgcggtt cctccgacct gtacctggtt 360
283 acccgtcacg ctgacgttat cccggttcgt cgtcgtggtg actcccgtgg ttccctgctg 420
285 tccccgcgtc cgatctccta cctgaaaggt tcctccggtg tccgctgct gtgcccggct 480
287 ggtcacgctg ttggtatctt ccgtgctgct gtttgcaccc gtggtgttgc taaagctgtt 540
289 gacttcatcc cggttgaatc cctggaaacc accatgcgtt ccccgtga             588
293 <210> SEQ ID NO: 5
295 <211> LENGTH: 15
297 <212> TYPE: PRT
299 <213> ORGANISM: Hepatitis C virus
303 <400> SEQUENCE: 5
305 Gln Gln Thr Arg Gly Leu Leu Gly Cys Ile Ile Thr Ser Leu Thr
307  1               5                  10                  15
313 <210> SEQ ID NO: 6
315 <211> LENGTH: 15
317 <212> TYPE: PRT
319 <213> ORGANISM: Hepatitis C virus
323 <400> SEQUENCE: 6
325 Gln Gln Thr Arg Gly Glu Glu Gly Cys Gln Glu Thr Ser Gln Thr
327  1               5                  10                  15
333 <210> SEQ ID NO: 7
335 <211> LENGTH: 15
337 <212> TYPE: PRT
339 <213> ORGANISM: Hepatitis C virus
343 <400> SEQUENCE: 7
345 Gln Gln Thr Arg Gly Glu Glu Gly Cys Gln Gln Thr Ser Glu Thr
347  1               5                  10                  15
353 <210> SEQ ID NO: 8
355 <211> LENGTH: 15
357 <212> TYPE: PRT
359 <213> ORGANISM: Hepatitis C virus
363 <400> SEQUENCE: 8
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367  1               5                  10                  15
373 <210> SEQ ID NO: 9
375 <211> LENGTH: 15
377 <212> TYPE: PRT
379 <213> ORGANISM: Hepatitis C virus
383 <400> SEQUENCE: 9
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387  1               5                  10                  15
393 <210> SEQ ID NO: 10
395 <211> LENGTH: 15
397 <212> TYPE: PRT
399 <213> ORGANISM: Hepatitis C virus
403 <400> SEQUENCE: 10
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413 <210> SEQ ID NO: 11
415 <211> LENGTH: 15
417 <212> TYPE: PRT
419 <213> ORGANISM: Hepatitis C virus
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427   1             5                   10                      15
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435 <211> LENGTH: 195
437 <212> TYPE: PRT
439 <213> ORGANISM: Hepatitis C virus
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447   1             5                   10                      15
451 Gly Ala Tyr Ala Gln Gln Thr Arg Gly Glu Glu Gly Cys Gln Glu Thr
453              20                  25                  30
457 Ser Gln Thr Gly Arg Asp Lys Asn Gln Val Glu Gly Glu Val Gln Ile
459             35                  40                  45
463 Val Ser Thr Ala Ala Gln Thr Phe Leu Ala Thr Cys Ile Asn Gly Val
465        50                  55                  60
469 Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr Ile Ala Ser Pro
471  65                  70                  75                  80
475 Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp Lys Asp Leu Val
477                  85                  90                  95
481 Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu Thr Pro Cys Thr Cys
483              100                 105                 110
487 Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala Asp Val Ile Pro
489          115                 120                 125
493 Val Arg Arg Arg Gly Asp Ser Arg Gly Ser Leu Leu Ser Pro Arg Pro
495        130                 135                 140
499 Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu Leu Cys Pro Ala
501 145                 150                 155                 160
505 Gly His Ala Val Gly Ile Phe Arg Ala Ala Val Cys Thr Arg Gly Val
507                  165                 170                 175
511 Ala Lys Ala Val Asp Phe Ile Pro Val Glu Ser Leu Glu Thr Thr Met
513              180                 185                 190
517 Arg Ser Pro
519          195
525 <210> SEQ ID NO: 13
527 <211> LENGTH: 588
529 <212> TYPE: DNA
531 <213> ORGANISM: Hepatitis C virus
535 <400> SEQUENCE: 13
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539 cagcagactc gaggtgagga gggttgccaa gaaacctccc agaccggtcg tgacaaaaac 120
541 caggttgaag gtgaagttca gatcgtttcc accgctgctc agaccttcct ggctacctgc 180
543 atcaacggtg tttgctggac cgtttaccac ggtgctggta cccgtaccat cgcttccccg 240
545 aaaggtccgg ttatccagat gtacaccaac gttgacaaag acctggttgg ttggccggct 300
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547 ccgcagggtt cccgttccct gaccccgtgc acctgcggtt cctccgacct gtacctggtt 360
549 acccgtcacg ctgacgttat cccggttcgt cgtcgtggtg actcccgtgg ttccctgctg 420
551 tccccgcgtc cgatctccta cctgaaaggt tcctccggtg gtccgctgct gtgcccggct 480
553 ggtcacgctg ttggtatctt ccgtgctgct gtttgcaccc gtggtgttgc taaagctgtt 540
555 gacttcatcc cggttgaatc cctggaaacc accatgcgtt ccccgtga           588
559 <210> SEQ ID NO: 14
561 <211> LENGTH: 197
563 <212> TYPE: PRT
565 <213> ORGANISM: Hepatitis C virus
569 <400> SEQUENCE: 14
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573  1               5                  10                  15
577 Gly Asp Thr Ala Tyr Ala Gln Gln Thr Arg Gly Glu Glu Gly Cys Gln
579              20                  25                  30
583 Glu Thr Ser Gln Thr Gly Arg Asp Lys Asn Gln Val Glu Gly Glu Val
585              35                  40                  45
589 Gln Ile Val Ser Thr Ala Ala Gln Thr Phe Leu Ala Thr Cys Ile Asn
591      50                  55                  60
595 Gly Val Cys Trp Thr Val Tyr His Gly Ala Gly Thr Arg Thr Ile Ala
597  65                  70                  75                  80
601 Ser Pro Lys Gly Pro Val Ile Gln Met Tyr Thr Asn Val Asp Lys Asp
603              85                  90                  95
607 Leu Val Gly Trp Pro Ala Pro Gln Gly Ser Arg Ser Leu Thr Pro Cys
609              100                 105                 110
613 Thr Cys Gly Ser Ser Asp Leu Tyr Leu Val Thr Arg His Ala Asp Val
615          115                 120                 125
619 Ile Pro Val Arg Arg Arg Gly Asp Ser Arg Gly Ser Leu Leu Ser Pro
621      130                 135                 140
625 Arg Pro Ile Ser Tyr Leu Lys Gly Ser Ser Gly Gly Pro Leu Leu Cys
627 145                 150                 155                 160
631 Pro Ala Gly His Ala Val Gly Ile Phe Arg Ala Ala Val Cys Thr Arg
633              165                 170                 175
637 Gly Val Ala Lys Ala Val Asp Phe Ile Pro Val Glu Ser Leu Glu Thr
639              180                 185                 190
643 Thr Met Arg Ser Pro
645          195
651 <210> SEQ ID NO: 15
653 <211> LENGTH: 594
655 <212> TYPE: DNA
657 <213> ORGANISM: Hepatitis C virus
661 <400> SEQUENCE: 15
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665 tacgctcagc agactcgagg tgaggagggt tgccaagaaa cctcccagac cggtcgtgac 120
667 aaaaaccagg ttgaaggtga agttcagatc gtttccaccg ctgctcagac cttcctggct 180
669 acctgcatca cggtgtttg ctggaccgtt taccacggtg ctggtacccg taccatcgct 240
671 tccccgaaag gtccggttat ccagatgtac accaacgttg acaaagacct ggttggttgg 300
673 ccggctccgc agggttcccg ttccctgacc ccgtgcacct gcggttcctc cgacctgtac 360
675 ctggttaccc gtcacgctga cgttatcccg gttcgtcgtc gtggtgactc ccgtggttcc 420
677 ctgctgtccc cgcgtccgat ctcctacctg aaaggttcct ccggtggtcc gctgctgtgc 480
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VERIFICATION SUMMARY   DATE: 10/25/2001
PATENT APPLICATION: US/09/965,594   TIME: 08:46:15

Input Set : **A:\PTO.AMC.txt**
Output Set: **N:\CRF3\10252001\I965594.raw**

L:27 M:270 C: Current Application Number differs, Replaced Current Application Number
L:29 M:271 C: Current Filing Date differs, Replaced Current Filing Date